

## WEST Search History

DATE: Thursday, August 09, 2007

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	<i>DB=PGPB,USPT,USOC,EPAB,JPAB,DWPI; PLUR=YES; OP=OR</i>		
<input type="checkbox"/>	L24	L20 and (clostrid\$ or tetan\$)	62
<input type="checkbox"/>	L23	L20 and (tetan\$)	36
<input type="checkbox"/>	L22	L20 and (clostrid\$ or tetan\$)	62
<input type="checkbox"/>	L21	L20 and (clostrid\$ or fragment or tetan\$)	269
<input type="checkbox"/>	L20	l14 and \$toxin.clm. not l16 not l17	401
<input type="checkbox"/>	L19	l15 and \$toxin not l16 not l17	0
<input type="checkbox"/>	L18	l15 and \$toxin.clm. not l16 not l17	0
<input type="checkbox"/>	L17	l15 not l16	26
<input type="checkbox"/>	L16	L14 same tetan\$.clm.	22
<input type="checkbox"/>	L15	L14 and tetan\$.clm.	48
<input type="checkbox"/>	L14	(\$spor\$ or sporu\$ or \$spore).clm.	147648
	<i>DB=PGPB; PLUR=YES; OP=OR</i>		
<input type="checkbox"/>	L13	US-20050232947-A1.did.	1
	<i>DB=PGPB,USPT,USOC,EPAB,JPAB,DWPI; PLUR=YES; OP=OR</i>		
<input type="checkbox"/>	L12	L11 and tetan\$	19
<input type="checkbox"/>	L11	(\$spor\$ or sporu\$ or \$spore).ti.	4280
<input type="checkbox"/>	L10	L9 same (foreign or heterologous or heter-ologous or recombinant\$ or plasmid or vector or cassette or (delivery near2 system)).clm.	53
<input type="checkbox"/>	L9	tetanus!.clm. same \$toxin.clm.	306
<input type="checkbox"/>	L8	l2 and \$toxin.clm.	64
<input type="checkbox"/>	L7	l2 and (tetanus or fragment-c or c-fragment or Hc or bindingdomain or cellbinding).clm.	0
<input type="checkbox"/>	L6	l2 and (tetanus or fragment-c or c-fragment or heavy or Hc or bindingdomain or cellbinding).clm.	13
<input type="checkbox"/>	L5	L4 and (foreign or heterologous or heter-ologous or recombinant\$ or plasmid or vector or cassette or (delivery near2 system))	89
<input type="checkbox"/>	L4	(rrno or rrn) near3 (p1 or p2 or operon)	93
<input type="checkbox"/>	L3	L2 and (\$spor\$ or sporu\$ or \$spore).clm.	38
<input type="checkbox"/>	L2	host.clm. same bacill\$.clm.	759
<input type="checkbox"/>	L1	tetanus! near10 (spore\$ or \$spore)	48

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L26: Entry 9 of 30

File: PGPB

May 8, 2003

DOCUMENT-IDENTIFIER: US 20030088081 A1

TITLE: High level expression of immunogenic proteins in the plastids of higher plants

Detail Description Paragraph:

[0120] The map of the tobacco plastid lox >aadA> deletion vector is shown in FIG. 15. It is the basic lox deletion vector with an aadA gene cloned into the BglII sites oriented towards the rrn operon.

Detail Description Paragraph:

[0201] Plasmids pJST10 and pJST11 were obtained by replacing the neo coding region in plasmid pHK40 with the tetC-AT and tetC-GC coding regions as an NdeI-XbaI fragments. Plasmid pHK40 is a plastid transformation vector derived from plasmid pPRV111A (Kuroda and Maliga 2001a) with a spectinomycin resistance (aadA) gene as a selective marker and a neo gene expressed in a cassette consisting of a PrnLT7g10 cassette and the rbcL 3'-UTR (TrbcL). The tetC genes are divergently oriented relative to the rrn operon (FIG. 21B).

Detail Description Paragraph:

[0202] Plasmid pJST12 was obtained by replacing the neo coding region in plasmid pHK73 with the tetC-AT coding region as an NdeI-XbaI fragment (SEQ. ID No: 31). Plastid transformation vector pHK73 is a pPRV111B vector derivative in which the neo coding region is expressed in a cassette consisting of a PrnLatpB cassette (plastid rrn operon promoter fused with atpB leader and an NdeI site including the ATG) and TrbcL. The PrnLatpB cassette is identical with the cassette in plasmid pHK10 (Kuroda and Maliga 2001b), except that an NdeI site was created by replacing AT with a CA at the -3/-2 position upstream of the ATG. The tetC gene in plasmid pJST12 is in tandem orientation with the rrn operon (FIG. 21B).

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L29: Entry 27 of 117

File: USPT

Apr 13, 2004

DOCUMENT-IDENTIFIER: US 6720167 B1

TITLE: Insecticidal bacteria, and methods for making and using them

Detailed Description Text (72):

Agaisse 1996 reviewed databases and identified numerous examples of putative STAB-SD sequences in 5' untranslated regions ("UTR"), including those of four cry genes from Bt, the cwp locus of B. brevis, and the inIAB locus of Listeria monocytogenes. Any of these STAB-SD sequences can be used to produce high levels of Bs toxin in Bacillus when placed between a strong Bacillus promoter and a ribosome binding site. The STAB-SD sequences identified share fairly high homology to one another.

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